

results of search on Crystallography Journals Online

20 articles match your search Aurora-A kinase.

[modify](#) your search.

Select bibliographic records for downloading using the checkboxes or **select all** button

Results 11 to 20, sorted by relevance:

[Sort by journal](#)

[Sort by year](#)

[Concise format](#)

[Previous results](#)

[Select all 20 hits](#)



[pdf](#)

[similar papers](#)

[open access](#)

Acta Cryst. (2007). A63, s127

Bone morphogenetic protein type II receptor structure in two crystal forms

S. Cutfield, J. Cutfield and P. Mace



[pdf](#)

[similar papers](#)

[open access](#)

Acta Cryst. (2007). A63, s127

Interaction of SRP54 GTPase and SRP RNA in the free signal recognition particle

T. Hainzl, S. Huang and A. E. Sauer-Eriksson



[pdf](#)

[similar papers](#)

[open access](#)

Acta Cryst. (2008). A64, C36-C37

A new beamline to achieve protein micro-crystallography at SPring-8

K. Hirata, A. Nisawa, G. Ueno, N. Shimizu, T. Kumasaka, T. Tanaka, S. Takahashi, K. Takeshita, H. Ohashi, S. Goto, H. Kitamura and M. Yamamoto



[pdf](#)

[similar papers](#)

[open access](#)

Acta Cryst. (2008). A64, C37

Microcrystallography at Diamond: facilities for crystal optimization and structure determination

G. Evans, A. Wagner, E. Shepherd, J. Aishima, M. Burt, K. Wilkinson, M. Harding, A. Taylor, S. Green, G. McIntyre and R. Holdsworth



[pdf](#) [similar papers](#) [open access](#)

Acta Cryst. (2008). **A64**, C37

Fragment-based drug discovery: from crystal to clinic

J. Patel



[pdf](#) [similar papers](#) [open access](#)

Acta Cryst. (2008). **A64**, C37

Role of structures in designing anti-AIDS drugs targeting reverse transcriptase

K. Das, J. D. Baruman, R. Bandwar, A. D. Clark Jr, S. H. Hughes and E. Arnold



[pdf](#) [similar papers](#) [open access](#)

Acta Cryst. (2008). **A64**, C37-C38

Monoamine oxidases and LSD1: similar chemistry for neurotransmitter and chromatin modification

A. Mattevi, F. Forneris, D. E. Edmondson and C. Binda



[pdf](#) [similar papers](#) [open access](#)

Acta Cryst. (2007). **A63**, s128

Structural basis for mRNA degradation by the RnaseJ

I. Li de la Sierra-Gallay, L. Zig, A. Jamalli and H. Putzer



[pdf](#) [similar papers](#) [open access](#)

Acta Cryst. (2007). **A63**, s128-s129

Unexpected domain architecture of Type IIP restriction endonuclease SdaI

G. Tamulaitiene, S. Grazulis and V. Siksnyš

[pdf](#)[similar papers](#)[open access](#)

Acta Cryst. (2007). **A63**, s128

Structural analysis of SmeT, a repressor of the *S. maltophilia* multidrug efflux pump SmeDEF

M. J. Mate-Perez, A. Hernandez, J. L. Martinez and A. Romero

[Sort by journal](#)[Sort by year](#)[Concise format](#)[Previous results](#)[Select all 20 hits](#)[download](#)selection in [BIBTeX](#)

format

List from result number: 11

Find reference:

Volume

Page

Search:

From

to

[Advanced search](#)